

			ATTY. DOCKET NO. 06137.0021.US02	APPLICATION NO. 09/181,601																																																																																									
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			FILING DATE: October 29, 1998	GROUP ART UNIT: 1643/655
OTHER (Including Author, Title, Date, Pertinent Pages, etc.)				
<i>M</i>	AA	<u>4</u> ✓	Yamazaki, <i>et al.</i> , "Secondary structure and signal assignments of human-immunodeficiency-virus-1 protease complexed to a novel, structure-based inhibitor" <i>Eur. J. Biochem.</i> 219 : 707-12 (1994). ✓	
<i>L</i>	AB	<u>4</u> ✓	Zimmerman, <i>et al.</i> , "Automated sequencing of amino acid spin systems in proteins using multidimensional HCC(CO)NH-TOCSY spectroscopy and constraint propagation methods from artificial intelligence" <i>J. Biomol. NMR</i> 4 : 241-256 (1994). ✗	
<i>L</i>	AC	<u>4</u> ✓	Zimmerman, <i>et al.</i> , "Automated analysis of nuclear magnetic resonance assignments for proteins" <i>Curr. Opin. Struct. Bio.</i> 5 : 644-73 (1995). ✗	
<i>M</i>	AD	<u>4</u> ✓	Zimmerman, <i>et al.</i> , "Automated Analysis of Protein NMR Assignments Using Methods from Artificial Intelligence" <i>J. Mol. Bio.</i> 269 : 592-610 (1997). ✗	
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OTHER (Including Author, Title, Date, Pertinent Pages, etc.)

<i>M</i>	AA	1 ✓	Bagby, <i>J. Biomol.</i> , "The button test: A small scale method using microdialysis cells for assessing protein solubility at concentrations suitable for NMR" <i>NMR</i> 10 : 279-82 (1997).
	AB	1 ✓	Billeter, <i>et al.</i> , "Sequential Resonance Assignments in Protein ^1H Nuclear Magnetic Resonance Spectra" <i>J. Mol. Biol.</i> 155 : 321-46 (1982).
	AC	1 ✓	Billeter, <i>et al.</i> , "A Program for Semi-automatic Sequential Resonance Assignments in Protein ^1H Nuclear Magnetic Resonance Spectra" <i>J. Magn. Resonance</i> 76 : 400-415 (1988).
	AD	1 ✓	Bork, <i>et al.</i> , "From genome sequences to protein function" <i>Curr. Opin. Struct. Bio.</i> 4 : 393-403 (1994).
	AE	1 ✓	Branden, <i>et al.</i> <u>Introduction to Protein Structure</u> pp. 11-31 and 43-57 (New York: Garland Publishing, Inc., 1991).
	AF	1 ✓	Brenner, <i>et al.</i> , "Assessing sequence comparison methods with reliable structurally identified distant evolutionary relationships" <i>Proc. Natl. Acad. Sci (U.S.A.)</i> 95 : 6073-78 (1998).
	AG	1 ✓	Chiang, <i>et al.</i> , "Structural Studies on the Cysteine-Rich N-terminal Domain of the Alzheimer's Amyloid β Peptide Precursor Protein" <i>Neurobiol. Aging, Supplement</i> 17(4S) : abstract 393 (1996).
	AH	1 ✓	Clore, <i>et al.</i> , "Multidimensional heteronuclear Nuclear Magnetic Resonance of Proteins" <i>Methods Enzymol.</i> 239 : 349-63 (1994).
	AI	1 ✓	Clore, <i>et al.</i> , "Structures of Larger Proteins, protein-Ligand and Protein-DNA Complexes by Multi-Dimensional Heteronuclear NMR" <i>Prog. Biophys. Mol. Bio.</i> 62 : 153-84 (1994).
	AJ	1 ✓	Clore, <i>et al.</i> , "Two-, Three-, and Four-Dimensional NMR Methods for Obtaining Larger and More Precise Three-Dimensional Structures of Proteins in Solution ¹ " <i>Ann. Rev. Biophys. Biophys. Chem.</i> 20 : 29-63 (1991).
	AK	1 ✓	Feng, <i>et al.</i> , "Solution NMR Structure and Backbone Dynamics of the Major Cold-Shock Protein (CspA) from <i>Escherichia coli</i> : Evidence for conformational Dynamics in the Single-Stranded RNA-Binding Site" <i>Biochem.</i> 37 : 10881-96 (1998).
	AL	1 ✓	Friedrichs, <i>et al.</i> , An automated procedure for the assignment of protein $^1\text{H}\text{N}$, ^{15}N , $^{13}\text{C}^\alpha$, $^1\text{H}^\alpha$, $^{13}\text{C}^\beta$ and $^1\text{H}^\beta$ resonances" <i>J. Biomol. NMR</i> 4 : 703-726 (1994).
	AM	1 ✓	Gibrat, <i>et al.</i> , "Surprising similarities in structure comparison" <i>Current Opinion in Structural Biology</i> 6 : 377-85 (1996).
<i>M</i>	AN	1 ✓	Go, "Protein structures and split genes" <u>Advances in Biophysics</u> 19 : 91-131 (Elsevier, 1995).
<i>M</i>	AO	1 ✓	Gouzey, <i>et al.</i> , "Graphical interface for ProDom domain families" <i>Trends Biochem. Sci.</i> , 21 : 493 (1994).

EXAMINER

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<i>O I P E</i> JUL 01 1999 PATENT & TRADEMARK OFFICE	AA	3 ✓	Montelione, <i>et al.</i> , "Solution Structure of Murine Epidermal Growth Factor Determined by NMR Spectroscopy and Refined by Energy Minimization with Restraints" <i>Biochemistry</i> 31 : 236-49 (1992).	
	AB	3 ✓	Mumenthaler, <i>et al.</i> , "Automated Assignment of Simulated and Experimental NOESY Spectra of Proteins by Feedback Filtering and Self-correcting Distance Geometry" <i>J. Mol. Biol.</i> 254 : 465-480 (1995).	
	AC	3 ✓	Nilges, "Calculation of Protein Structures with Ambiguous Distance Restraints. Automated Assignment of Ambiguous NOE Crosspeaks and Disulphide Connectivities" <i>J. Mol. Biol.</i> 245 : 645-60 (1995).	
	AD	3 ✓	Olsen, <i>et al.</i> , "Evaluation of an algorithm for the automated sequential assignment of protein backbone resonances: A demonstration of the connectivity tracing assignment tools (CONTRAST) software package" <i>J. Biomol. NMR</i> 4 : 385-410 (1994).	
	AE	3 ✓	Orengo, "CATH – a hierachic classification of protein domain structures" <i>Structure</i> 5 : 1093-1108 (1997).	
	AF	3 ✓	Patthy, <i>et al.</i> , "Introns and Exons" <i>Curr. Opin. Struc. Biol.</i> 4 : 383-92 (1994).	
	AG	3 ✓	Rios, <i>et al.</i> , "Phase labeling of C-H and C-C spin-system topologies: Application in constant-time PFG-CBCA(CO)NH experiments for discriminating amino acid spin-system types" <i>J. Biomol. NMR</i> 8 : 345-350 (1996).	
	AH	3 ✓	Sandbrink, <i>et al.</i> , "Alternative Splicing Generates Functionally Related isoforms ^a " <i>Ann. NY Acad. Sci.</i> 777 : 281-87 (1996).	
	AI	3 ✓	Shang, <i>et al.</i> , "Sensitivity Enhancement of Triple-Resonance Protein NMR Spectra by Proton Evolution of Multiple-Quantum Coherences using a Simultaneous ¹ H and ¹³ C Constant-Time Evolution Period" <i>J. Amer. Chem. Soc.</i> 119 : 9274-78 (1997).	
	AJ	3 ✓	Shimotakahara, <i>et al.</i> , "NMR Structural Analysis of an Analog of an Intermediate Formed in the Rate-Determining Step of One Pathway in the Oxidative Folding of Bovine Pancreatic Ribonuclease A: Automated Analysis of ¹ H, ¹³ C, and ¹⁵ N Resonance Assignments for Wild-Type and [C65S, C72S] Mutant Forms" <i>Biochem.</i> 36 : 6915-29 (1997).	
	AK	3 ✓	Swapana, <i>et al.</i> , "Application of multiple-quantum line narrowing with simultaneous ¹ H and ¹³ C constant-time scalar-coupling evolution in PFG-HACANH and PFG-HACA(CO)NH triple-resonance experiments" <i>J. Biomol. NMR</i> 9 : 105-111 (1997).	
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	AM	3 ✓	Wuthrich, "Protein Structure Determination in Solution by Nuclear Magnetic Resonance Spectroscopy" <i>Science</i> 243 : 45-50 (1989).	
	AN	3 ✓	Wuthrich, "NMR – This Other Method for Protein and Nucleic Acid Structure Determination" <i>Acta Cryst. Section D</i> 51 : 249-70 (1995).	
	AO	3 ✓	Yashikai, <i>et al.</i> , "Genomic organization of the human amyloid beta-protein precursor gene" <i>Gene</i> 87 : 257-63 (1990).	
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